73.   The vector according to claim 72, wherein said nucleotide sequence encoding an immunoglobulin light chain variable region is selected from the group consisting of Vκ1 (SEQ ID NO: 42), Vκ2 (SEQ ID NO: 44), Vκ3 (SEQ ID NO: 46), Vκ4 (SEQ ID NO: 48), Vλ1 (SEQ ID NO: 50), Vλ2 (SEQ ID NO: 52), and Vλ3 (SEQ ID NO: 54).

74.   The vector according to claim 71, wherein said nucleotide sequence encoding a (poly)peptide encodes an immunoglobulin heavy chain variable region comprising a modular sequence of four consensus framework regions interspaced by three complementary determining regions CDR1, CDR2, and CDR3,

wherein said nucleotide sequence comprises DNA cleavage sites at the boundary between each consensus framework region and each complementary determining region, and wherein each of said cleavage sites is unique within said vector.

75.   The vector according to claim 74, wherein said nucleotide sequence encoding an immunoglobulin heavy chain variable region is selected from the group consisting of VH1A (SEQ ID NO: 56), VH1B (SEQ ID NO: 58), VH2 (SEQ ID NO: 60), VH3 (SEQ ID NO: 62), VH4 (SEQ ID NO: 64), VH5 (SEQ ID NO: 66), and VH6 (SEQ ID NO: 68).

76.   The vector according to claim 71, wherein said nucleotide encoding a (poly)peptide encodes an immunoglobulin constant chain region selected from the group consisting of Cκ (SEQ ID NO: 165), Cλ (SEQ ID NO: 169), and CH1 (SEQ ID NO: 167).

77.   The vector according to claim 66, wherein said nucleotide sequence encoding a (poly)peptide encodes a (poly)peptide selected from the group consisting of an immunoglobulin light chain variable region, an immunoglobulin heavy chain variable region, and an immunoglobulin constant chain region.

78.   The vector according to claim 77, wherein said nucleotide sequence encoding a (poly)peptide encodes an immunoglobulin light chain variable region comprising a modular sequence of four consensus framework regions interspaced by three complementary determining regions CDR1, CDR2, and CDR3,

wherein said nucleotide sequence comprises DNA cleavage sites at the boundary between each consensus framework region and each complementary determining region, and wherein each of said cleavage sites is unique within said vector.

79.   The vector according to claim 78, wherein said nucleotide sequence encoding an immunoglobulin light chain variable region is selected from the group consisting of Vκ1 (SEQ ID NO: 42), Vκ2 (SEQ ID NO: 44), Vκ3 (SEQ ID NO: 46), Vκ4 (SEQ ID NO: 48), Vλ1 (SEQ ID NO: 50), Vλ2 (SEQ ID NO: 52), and Vλ3 (SEQ ID NO: 54).

80.   The vector according to claim 77, wherein said nucleotide sequence encoding a (poly)peptide encodes an immunoglobulin heavy chain variable region comprising a modular sequence of four consensus framework regions interspaced by three complementary determining regions CDR1, CDR2, and CDR3,

wherein said nucleotide sequence comprises DNA cleavage sites at the boundary between each consensus framework region and each complementary determining region, and wherein each of said cleavage sites is unique within said vector.

81.   The vector according to claim 80, wherein said nucleotide sequence encoding an immunoglobulin heavy chain variable region is selected from the group consisting of VH1A (SEQ ID NO: 56), VH1B (SEQ ID NO: 58), VH2 (SEQ ID NO: 60), VH3 (SEQ ID NO: 62), VH4 (SEQ ID NO: 64), VH5 (SEQ ID NO: 66), and VH6 (SEQ ID NO: 68).

82.   The vector according to claim 77, wherein said nucleotide encodes poly)peptide encodes an immunoglobulin constant chain region selected from the group consisting of Cκ (SEQ ID NO: 165), Cλ (SEQ ID NO: 169), and CH1 (SEQ ID NO: 167).

83.   The vector according to claim 66, wherein said vector is a phagemid vector.--